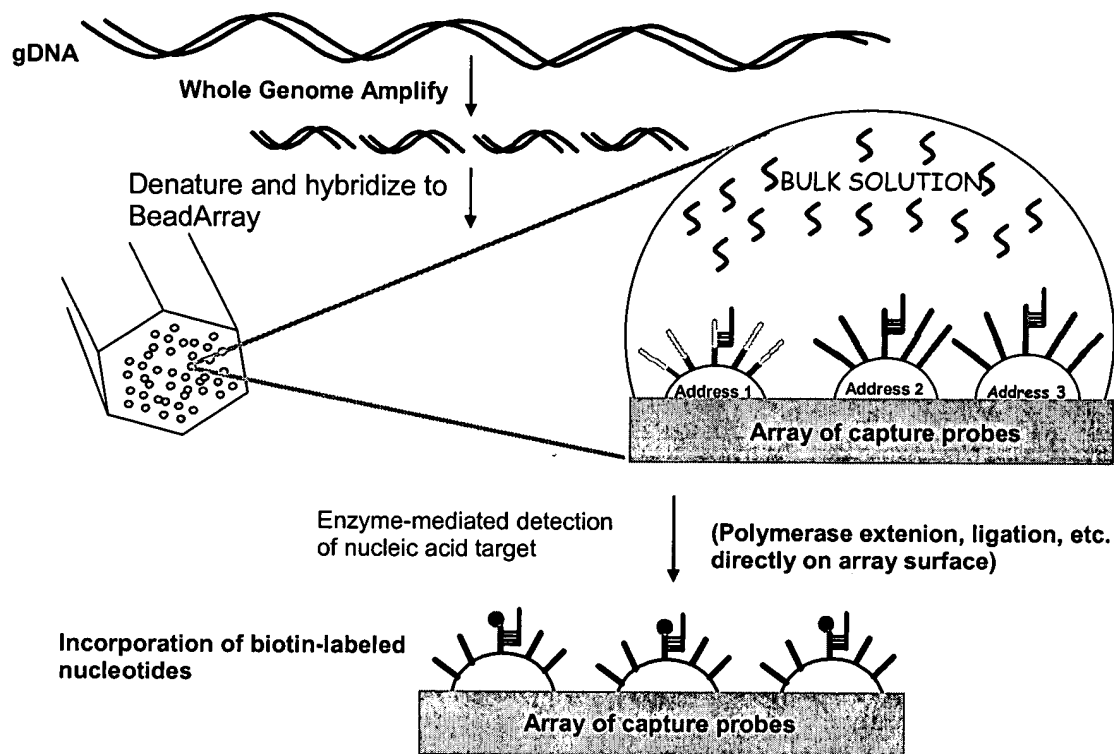
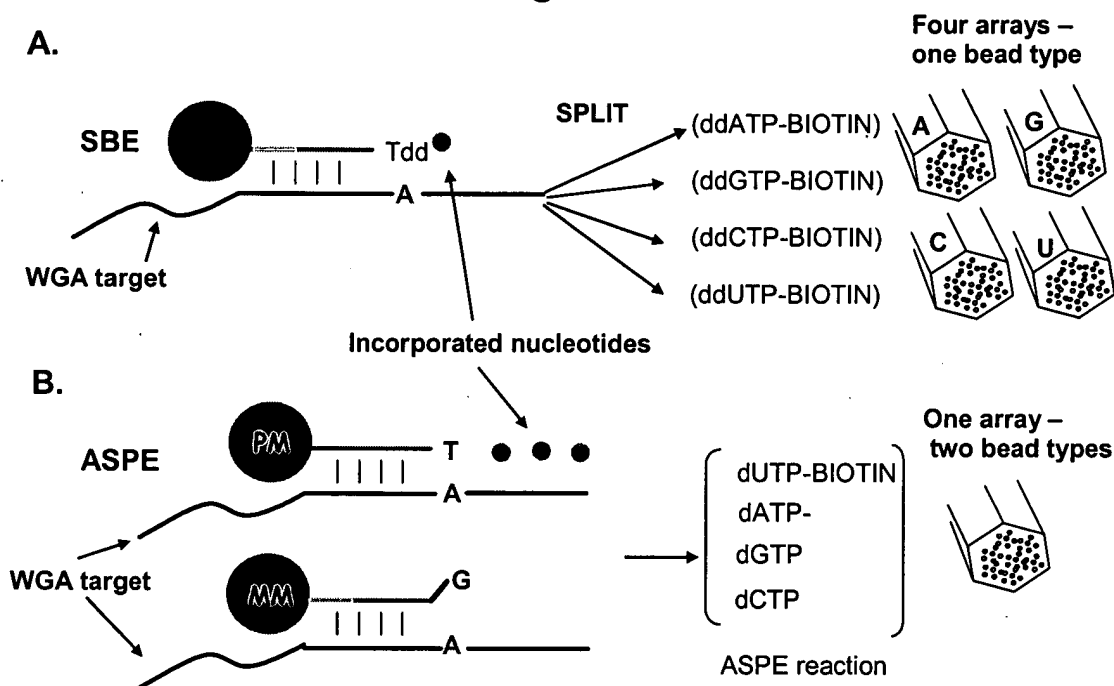


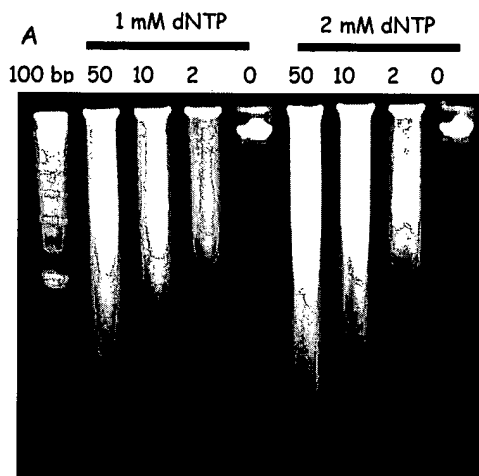
Figure 1



**Figure 2**



**Figure 3**



**B.**

Condition	dNTPs (mM)	Primer (uM)	Yield (ug)	Amplification
1	1	50	49.5	4950
2	1	10	28.5	2850
3	1	2	12.0	1200
4	1	0	6.0	600
5	2	50	79.5	7950
6	2	10	42.0	4200
7	2	2	21.0	2100
8	2	0	6.0	600

Figure 4

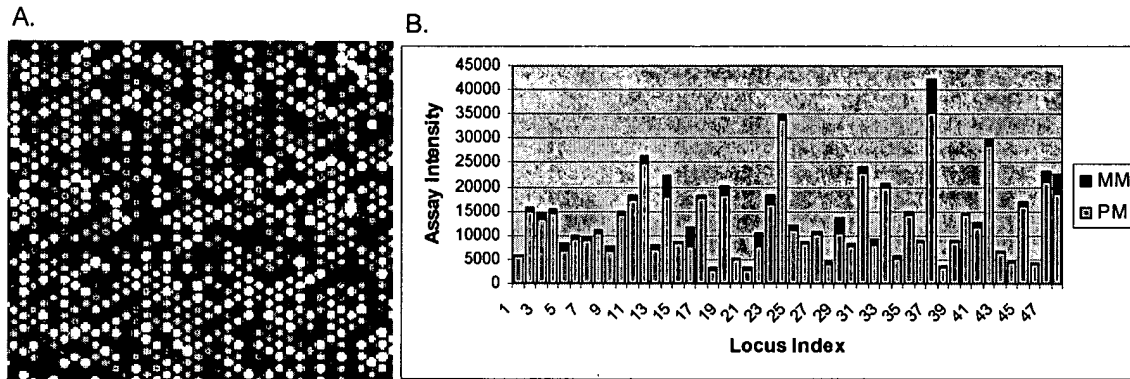


Figure 5

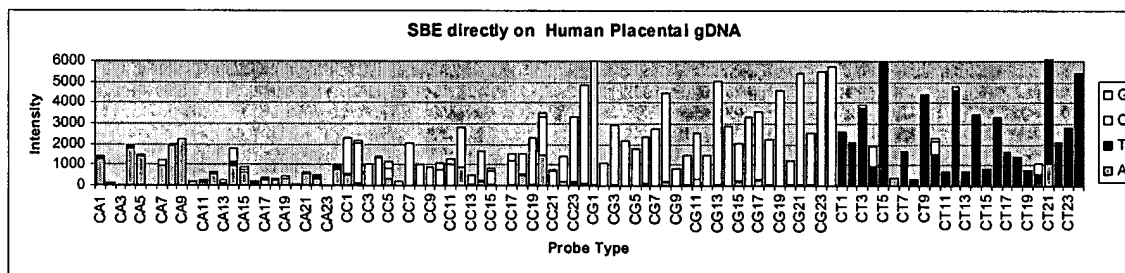


Figure 6

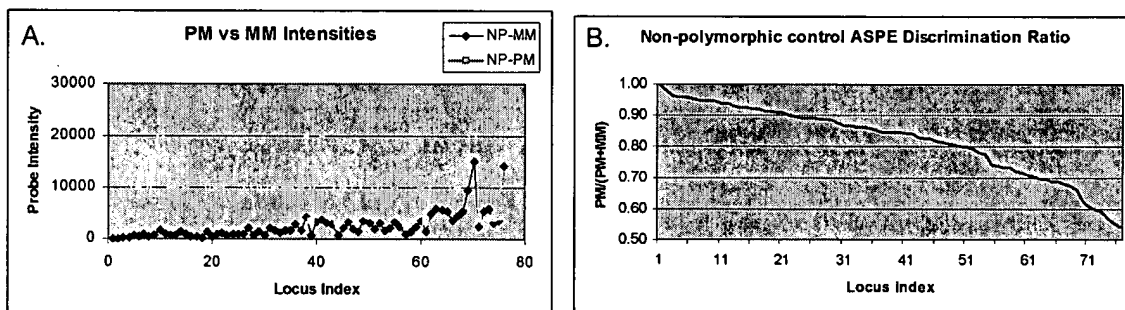


Figure 7

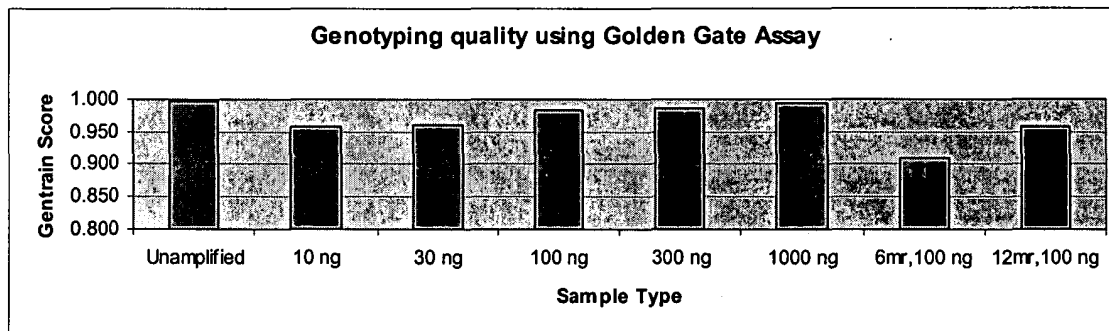
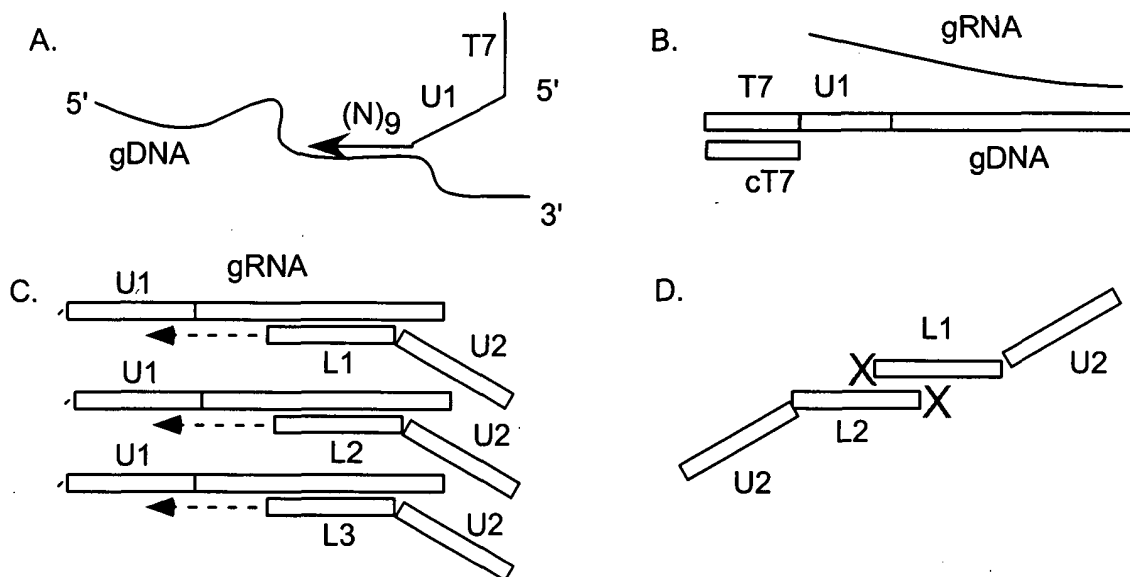
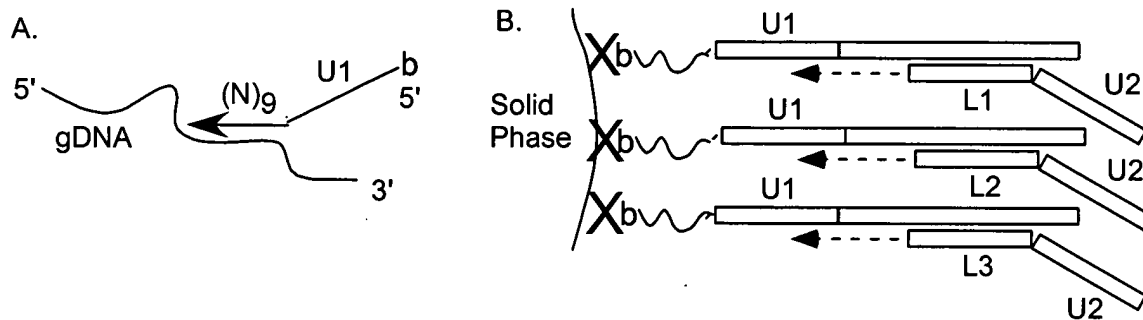


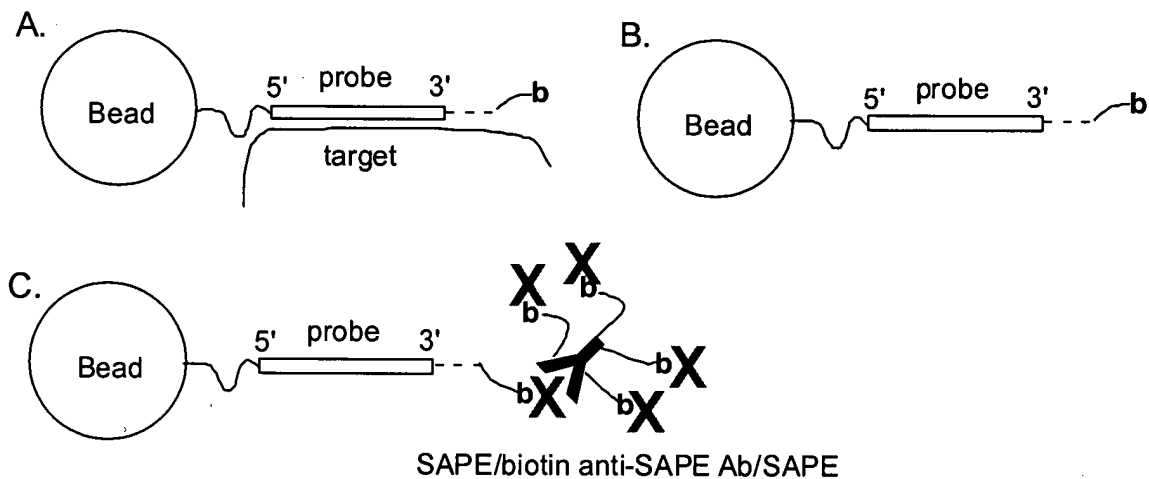
Figure 8



**Figure 9**

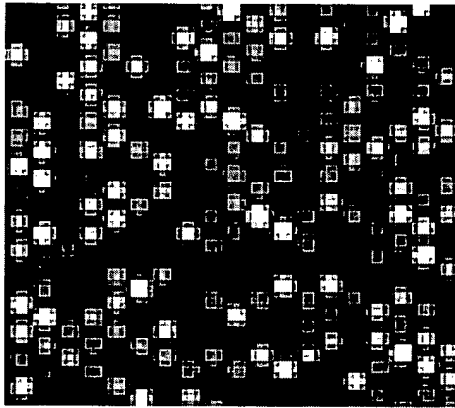


**Figure 10**

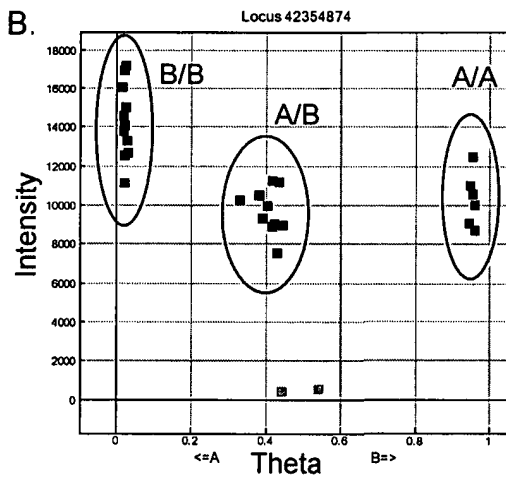


**Figure 11**

A.



B.



Title: METHODS AND COMPOSITIONS FOR WHOLE GENOME AMPLIFICATION  
AND GENOTYPING

Inventors: Gunderson et al.

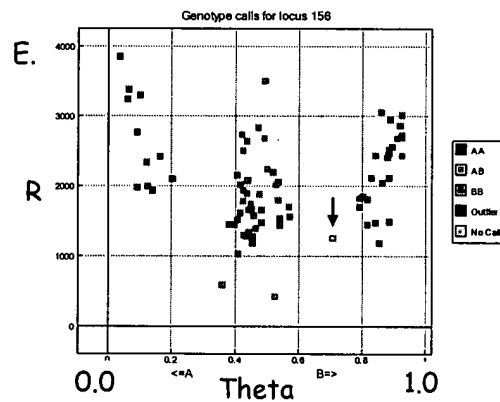
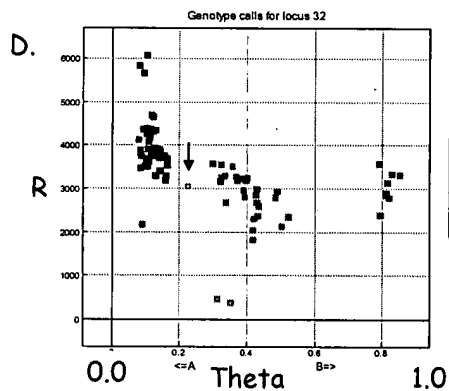
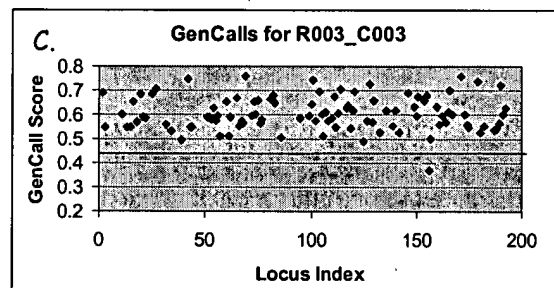
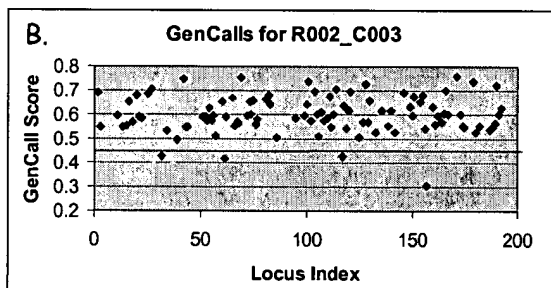
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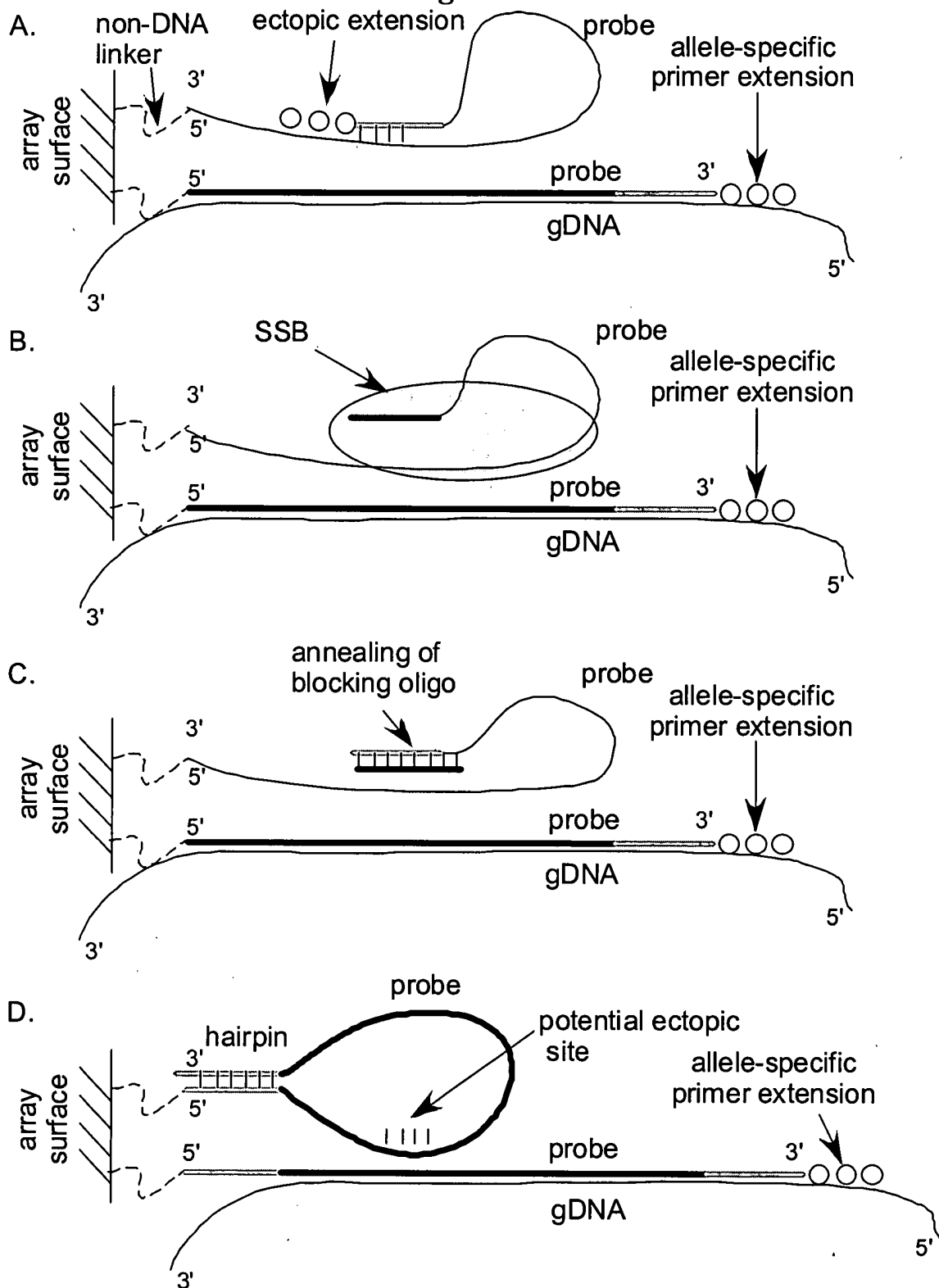
**Figure 12**

A.

Genotyping Accuracy:			
Call Rate	Accuracy	Calls	Correct Calls
100%	99.95%	4092	4090
99.70%	100%	4078	4078



**Figure 13**



**Figure 14**

